

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/581,468
Source: IFWP
Date Processed by STIC: 6/14/06

ENTERED



IFWP

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/581,468

DATE: 06/14/2006
TIME: 10:10:16

Input Set : A:\US2004-037600.txt
Output Set: N:\CRF4\06142006\J581468.raw

3 <110> APPLICANT: PDL BioPharma, Inc.
4 BALASA, Balaji
5 TSURUSHITA, Naoya
6 LANDOLFI, Nicholas F.
8 <120> TITLE OF INVENTION: TREATMENT OF INFLAMMATORY BOWEL DISEASES WITH ANTI-IP-10
9 ANTIBODIES
11 <130> FILE REFERENCE: 116 US PC01
C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/581,468
C--> 14 <141> CURRENT FILING DATE: 2006-06-01
16 <150> PRIOR APPLICATION NUMBER: PCT/US2004/014507
17 <151> PRIOR FILING DATE: 2004-05-07
19 <150> PRIOR APPLICATION NUMBER: US 60/527,882
20 <151> PRIOR FILING DATE: 2003-12-04
22 <160> NUMBER OF SEQ ID NOS: 79
24 <170> SOFTWARE: PatentIn version 3.3
26 <210> SEQ ID NO: 1
27 <211> LENGTH: 98
28 <212> TYPE: PRT
29 <213> ORGANISM: Homo sapiens
31 <400> SEQUENCE: 1
33 Met Asn Thr Ala Ile Leu Ile Cys Cys Leu Ile Phe Leu Thr Leu
34 1 5 10 15
37 Ser Gly Ile Gln Gly Val Pro Leu Ser Arg Thr Val Arg Cys Thr Cys
38 20 25 30
41 Ile Ser Ile Ser Asn Gln Pro Val Asn Pro Arg Ser Leu Glu Lys Leu
42 35 40 45
45 Glu Ile Ile Pro Ala Ser Gln Phe Cys Pro Arg Val Glu Ile Ile Ala
46 50 55 60
49 Thr Met Lys Lys Lys Gly Glu Lys Arg Cys Leu Asn Pro Glu Ser Lys
50 65 70 75 80
53 Ala Ile Lys Asn Leu Leu Lys Ala Val Ser Lys Glu Arg Ser Lys Arg
54 85 90 95
57 Ser Pro
61 <210> SEQ ID NO: 2
62 <211> LENGTH: 98
63 <212> TYPE: PRT
64 <213> ORGANISM: Homo sapiens
66 <400> SEQUENCE: 2
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69 1 5 10 15
72 Ser Gly Ile Gln Gly Val Pro Leu Ser Arg Thr Val Arg Cys Thr Cys
73 20 25 30
76 Ile Ser Ile Ser Asn Gln Pro Val Asn Pro Arg Ser Leu Glu Lys Leu

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77 35 40 45
 80 Glu Ile Ile Pro Ala Ser Gln Phe Cys Pro Arg Val Glu Ile Ile Ala
 81 50 55 60
 84 Thr Met Lys Lys Lys Gly Glu Lys Arg Cys Leu Asn Pro Glu Ser Lys
 85 65 70 75 80
 88 Ala Ile Lys Asn Leu Leu Lys Ala Val Ser Lys Glu Arg Ser Lys Arg
 89 85 90 95
 92 Ser Pro
 96 <210> SEQ ID NO: 3
 97 <211> LENGTH: 119
 98 <212> TYPE: PRT
 99 <213> ORGANISM: Mus sp.
 101 <400> SEQUENCE: 3
 103 Gln Ile Gln Leu Val Gln Ser Gly Pro Glu Leu Lys Lys Pro Gly Glu
 104 1 5 10 15
 107 Thr Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Asp Tyr
 108 20 25 30
 111 Ser Met His Trp Val Lys Gln Ala Pro Gly Lys Gly Leu Lys Trp Met
 112 35 40 45
 115 Gly Trp Ile Asn Thr Glu Ile Gly Glu Pro Thr Tyr Ala Asp Asp Phe
 116 50 55 60
 119 Lys Gly Arg Phe Ala Phe Ser Leu Glu Thr Ser Ala Ser Thr Ala Tyr
 120 65 70 75 80
 123 Leu Gln Ile Asn Asn Leu Lys Asn Glu Asp Thr Ala Thr Tyr Phe Cys
 124 85 90 95
 127 Ala Arg Asn Tyr Asp Tyr Asp Ala Tyr Phe Asp Val Trp Gly Ala Gly
 128 100 105 110
 131 Thr Thr Val Thr Val Ser Ser
 132 115
 135 <210> SEQ ID NO: 4
 136 <211> LENGTH: 107
 137 <212> TYPE: PRT
 138 <213> ORGANISM: Mus sp.
 140 <400> SEQUENCE: 4
 142 Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Leu Gly
 143 1 5 10 15
 146 Gly Lys Val Thr Ile Thr Cys Lys Ala Asp Gln Asp Ile Asn Lys Tyr
 147 20 25 30
 150 Ile Ala Trp Tyr Gln His Lys Pro Gly Arg Gly Pro Arg Leu Leu Leu
 151 35 40 45
 154 His His Thr Ser Thr Leu Gln Pro Gly Ile Pro Ser Arg Phe Ser Gly
 155 50 55 60
 158 Ser Gly Ser Gly Arg Asp Tyr Ser Phe Ser Ile Ser Asn Leu Glu Pro
 159 65 70 75 80
 162 Ala Asp Ile Ala Thr Tyr Tyr Cys Leu Gln Tyr Asp Ser Leu Leu Phe
 163 85 90 95
 166 Thr Phe Gly Ser Gly Thr Lys Leu Glu Ile Lys
 167 100 105
 170 <210> SEQ ID NO: 5

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171 <211> LENGTH: 5
172 <212> TYPE: PRT
173 <213> ORGANISM: Mus sp.
175 <400> SEQUENCE: 5
177 Asp Tyr Ser Met His
178 1 5
181 <210> SEQ ID NO: 6
182 <211> LENGTH: 17
183 <212> TYPE: PRT
184 <213> ORGANISM: Mus sp.
186 <400> SEQUENCE: 6
188 Trp Ile Asn Thr Glu Ile Gly Glu Pro Thr Tyr Ala Asp Asp Phe Lys
189 1 5 10 15
192 Gly
196 <210> SEQ ID NO: 7
197 <211> LENGTH: 10
198 <212> TYPE: PRT
199 <213> ORGANISM: Mus sp.
201 <400> SEQUENCE: 7
203 Asn Tyr Asp Tyr Asp Ala Tyr Phe Asp Val
204 1 5 10
207 <210> SEQ ID NO: 8
208 <211> LENGTH: 11
209 <212> TYPE: PRT
210 <213> ORGANISM: Mus sp.
212 <400> SEQUENCE: 8
214 Lys Ala Asp Gln Asp Ile Asn Lys Tyr Ile Ala
215 1 5 10
218 <210> SEQ ID NO: 9
219 <211> LENGTH: 7
220 <212> TYPE: PRT
221 <213> ORGANISM: Mus sp.
223 <400> SEQUENCE: 9
225 His Thr Ser Thr Leu Gln Pro
226 1 5
229 <210> SEQ ID NO: 10
230 <211> LENGTH: 9
231 <212> TYPE: PRT
232 <213> ORGANISM: Mus sp.
234 <400> SEQUENCE: 10
236 Leu Gln Tyr Asp Ser Leu Leu Phe Thr
237 1 5
240 <210> SEQ ID NO: 11
241 <211> LENGTH: 414
242 <212> TYPE: DNA
243 <213> ORGANISM: Mus sp.
245 <400> SEQUENCE: 11
246 atggcttggg tgtggacctt gctattcctg atggcagctg cccaaagtat ccaagcacag 60
248 atccagttgg tgcagtctgg acctgagctg aagaagcctg gagagacagt caagatctcc 120

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250	tgcaaggctt	ctggttatac	cttcacagac	tattcaatgc	actgggtgaa	gcaggctcca	180											
252	ggaaagggtt	taaagtggat	gggctggata	aacactgaga	ttggtgagcc	aacatatgca	240											
254	gatgacttca	agggacgggtt	tgccttctct	ttggaaacct	ctgccagcac	tgccctatttg	300											
256	cagatcaaca	acctcaaaaa	tgaggacacg	gctacatatt	tctgtctag	aaactatgat	360											
258	tacgacgcgt	acttcgatgt	ctggggcgca	gggaccacgg	tcaccgtctc	ctca	414											
261	<210>	SEQ ID NO:	12															
262	<211>	LENGTH:	381															
263	<212>	TYPE:	DNA															
264	<213>	ORGANISM:	Mus sp.															
266	<400>	SEQUENCE:	12															
267	atgagaccgt	ctattcagtt	cctggggctc	ttgttgttct	ggcttcatgg	tgctcagtgt	60											
269	gacatccaga	tgacacagtc	tccatcctca	ctgtctgcat	ctctggagg	caaagtccacc	120											
271	atcacttgca	aggcagacca	agacattaac	aagtatata	cttggtagcca	acacaaggcct	180											
273	ggaagaggtc	ctaggctgct	cctacatcac	acatctacat	tacagccagg	catccatca	240											
275	agttcagtg	gaagtgggtc	ttggagagat	tattcctca	gatcagcaa	cctggagcct	300											
277	gcagatatttgc	caacttatta	ttgtctacag	tatgatagtc	ttctattcac	gttcggctcg	360											
279	gggacaaagt	ttgaaataaaa	a				381											
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283	<211>	LENGTH:	119															
284	<212>	TYPE:	PRT															
285	<213>	ORGANISM:	Homo sapiens															
287	<400>	SEQUENCE:	13															
289	Glu	Val	Gln	Leu	Val	Gln	Ser	Gly	Ala	Glu	Val	Lys	Lys	Pro	Gly	Ala		
290	1				5					10					15			
293	Thr	Val	Lys	Ile	Ser	Cys	Lys	Val	Ser	Gly	Tyr	Thr	Phe	Thr	Asp	Tyr		
294					20				25					30				
297	Ser	Met	His	Trp	Val	Arg	Gln	Ala	Pro	Gly	Lys	Gly	Leu	Lys	Trp	Met		
298					35				40			45						
301	Gly	Trp	Ile	Asn	Thr	Glu	Ile	Gly	Glu	Pro	Thr	Tyr	Ala	Asp	Asp	Phe		
302					50				55			60						
305	Lys	Gly	Arg	Phe	Thr	Phe	Thr	Leu	Asp	Thr	Ser	Thr	Ser	Thr	Ala	Tyr		
306					65				70			75			80			
309	Met	Glu	Leu	Ser	Ser	Leu	Arg	Ser	Glu	Asp	Thr	Ala	Val	Tyr	Tyr	Cys		
310					85				90			95						
313	Ala	Arg	Asn	Tyr	Asp	Tyr	Asp	Ala	Tyr	Phe	Asp	Val	Trp	Gly	Gln	Gly		
314					100				105			110						
317	Thr	Thr	Val	Thr	Val	Ser	Ser											
318					115													
321	<210>	SEQ ID NO:	14															
322	<211>	LENGTH:	87															
323	<212>	TYPE:	PRT															
324	<213>	ORGANISM:	Homo sapiens															
326	<400>	SEQUENCE:	14															
328	Glu	Val	Gln	Leu	Val	Gln	Ser	Gly	Ala	Glu	Val	Lys	Lys	Pro	Gly	Ala		
329	1				5				10			15						
332	Thr	Val	Lys	Ile	Ser	Cys	Lys	Val	Ser	Gly	Tyr	Thr	Phe	Thr	Trp	Val		
333					20				25			30						
336	Gln	Gln	Ala	Pro	Gly	Lys	Gly	Leu	Glu	Trp	Met	Gly	Arg	Val	Thr	Ile		
337					35				40			45						

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Input Set : A:\US2004-037600.txt
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340 Thr Ala Asp Thr Ser Thr Asp Thr Ala Tyr Met Glu Leu Ser Ser Leu
341 50 55 60
344 Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys Ala Thr Trp Gly Gln Gly
345 65 70 75 80
348 Thr Thr Val Thr Val Ser Ser
349 85
352 <210> SEQ ID NO: 15
353 <211> LENGTH: 107
354 <212> TYPE: PRT
355 <213> ORGANISM: Homo sapiens
357 <400> SEQUENCE: 15
359 Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
360 1 5 10 15
363 Asp Arg Val Thr Ile Thr Cys Lys Ala Asp Gln Asp Ile Asn Lys Tyr
364 20 25 30
367 Ile Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Leu
368 35 40 45
371 His His Thr Ser Thr Leu Gln Pro Gly Ile Pro Ser Arg Phe Ser Gly
372 50 55 60
375 Ser Gly Ser Gly Arg Asp Tyr Thr Phe Thr Ile Ser Ser Leu Gln Pro
376 65 70 75 80
379 Glu Asp Ile Ala Thr Tyr Tyr Cys Leu Gln Tyr Asp Ser Leu Leu Phe
380 85 90 95
383 Thr Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys
384 100 105
387 <210> SEQ ID NO: 16
388 <211> LENGTH: 80
389 <212> TYPE: PRT
390 <213> ORGANISM: Homo sapiens
392 <400> SEQUENCE: 16
394 Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
395 1 5 10 15
398 Asp Arg Val Thr Ile Thr Cys Trp Tyr Gln Gln Lys Pro Gly Lys Ala
399 20 25 30
402 Pro Lys Leu Leu Ile Tyr Gly Val Pro Ser Arg Phe Ser Gly Ser Gly
403 35 40 45
406 Ser Gly Thr Asp Phe Thr Phe Thr Ile Ser Ser Leu Gln Pro Glu Asp
407 50 55 60
410 Ile Ala Thr Tyr Tyr Cys Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys
411 65 70 75 80
414 <210> SEQ ID NO: 17
415 <211> LENGTH: 412
416 <212> TYPE: DNA
417 <213> ORGANISM: Homo sapiens
419 <400> SEQUENCE: 17
420 acgcgtccac catgagacccg tctattcagt tcctggggct cttgttggtc tggcttcatg 60
422 gtgctcagtg tgacatccag atgacacagt ctccatcctc actgtctgca tctgtgggag 120
424 acagagtcac catcaactgc aaggcagacc aagacattaa caagtatata gcttggtacc 180
426 aacagaagcc tggaaaggct cctaagctgc tcctacatca cacatctaca ttacagccag 240

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 06/14/2006
PATENT APPLICATION: US/10/581,468 TIME: 10:10:18

Input Set : A:\US2004-037600.txt
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Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:21,22,23,24,25,26,27,28,29,30,31,32,33,34,35,36,37,38,39,40,51,52,53,54
Seq#:55,56,57,58,59,60,61,62,63,64,65,66,67,68,69,70,71,72

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/581,468

DATE: 06/14/2006

TIME: 10:10:18

Input Set : A:\US2004-037600.txt

Output Set: N:\CRF4\06142006\J581468.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application Number

L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date